

## SEQUENCE LISTING

<110> Genox Research, Inc.

National Center for Child Health and Development

<120> Methods for examination for allergic diseases, and  
drugs for treating allergic diseases

<130> G1-A0211-US

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<150> JP 2002-193841

<151> 2002-07-02

<160> 14

<170> PatentIn Ver. 2.0

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<212> DNA

<213> Homo sapiens

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Met Pro

tgt atc caa gcc caa tat ggg aca cca gca ccg agt ccg gga ccc cgt	164		
Cys Ile Gln Ala Gln Tyr Gly Thr Pro Ala Pro Ser Pro Gly Pro Arg			
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gac cac ctg gca agc gac ccc ctg acc cct gag ttc atc aag ccc acc	212		
Asp His Leu Ala Ser Asp Pro Leu Thr Pro Glu Phe Ile Lys Pro Thr			
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Met Asp Leu Ala Ser Pro Glu Ala Ala Pro Ala Ala Pro Thr Ala Leu			
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Pro Ser Phe Ser Thr Phe Met Asp Gly Tyr Thr Gly Glu Phe Asp Thr			
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Phe Leu Tyr Gln Leu Pro Gly Thr Val Gln Pro Cys Ser Ser Ala Ser			
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Ala Ser Phe Lys Phe Glu Asp Phe Gln Val Tyr Gly Cys Tyr Pro Gly			
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Tyr Tyr Gly Ser Pro Cys Ser Ala Pro Ser Pro Ser Thr Pro Ser Phe			
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Glu	Leu	Arg	Thr	Leu	Cys	Thr	Gln	Gly	Leu	Gln	Arg	Ile	Phe	Tyr	Leu
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Pro Gln Gly Ala Ser Pro Ala Ser Gln Ser Tyr Ser Tyr His Ser Ser  
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Gly Glu Tyr Ser Ser Asp Phe Leu Thr Pro Glu Phe Val Lys Phe Ser  
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Phe Ser Thr Phe Met Asp Asn Tyr Ser Thr Gly Tyr Asp Val Lys Pro			
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Pro Cys Leu Tyr Gln Met Pro Leu Ser Gly Gln Gln Ser Ser Ile Lys			
80	85	90	
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Val Glu Asp Ile Gln Met His Asn Tyr Gln Gln His Ser His Leu Pro			
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Pro Gln Ser Glu Glu Met Met Pro His Ser Gly Ser Val Tyr Tyr Lys			
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Pro Ser Ser Pro Pro Thr Pro Thr Thr Pro Gly Phe Gln Val Gln His			
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Ser Pro Met Trp Asp Asp Pro Gly Ser Leu His Asn Phe His Gln Asn			
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Tyr Val Ala Thr Thr His Met Ile Glu Gln Arg Lys Thr Pro Val Ser			
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Arg Leu Ser Leu Phe Ser Phe Lys Gln Ser Pro Pro Gly Thr Pro Val			
175	180	185	

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Lys Pro Lys Ser Pro Gln Glu Pro Ser Pro Pro Ser Pro Val Ser			
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Ala Asp Leu Pro Lys Ala Asp Gln Asp Leu Leu Phe Glu Ser Ala Phe			
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<211> 598

<212> PRT

<213> Homo sapiens

<400> 4

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Pro Ala Ser Gln Ser Tyr Ser Tyr His Ser Ser Gly Glu Tyr Ser Ser  
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Asp Phe Leu Thr Pro Glu Phe Val Lys Phe Ser Met Asp Leu Thr Asn  
 35 40 45

Thr Glu Ile Thr Ala Thr Thr Ser Leu Pro Ser Phe Ser Thr Phe Met  
 50 55 60

Asp Asn Tyr Ser Thr Gly Tyr Asp Val Lys Pro Pro Cys Leu Tyr Gln  
 65 70 75 80

Met Pro Leu Ser Gly Gln Gln Ser Ser Ile Lys Val Glu Asp Ile Gln  
 85 90 95

Met His Asn Tyr Gln Gln His Ser His Leu Pro Pro Gln Ser Glu Glu  
 100 105 110

Met Met Pro His Ser Gly Ser Val Tyr Tyr Lys Pro Ser Ser Pro Pro  
 115 120 125

Thr Pro Thr Thr Pro Gly Phe Gln Val Gln His Ser Pro Met Trp Asp

130 135 140

Asp Pro Gly Ser Leu His Asn Phe His Gln Asn Tyr Val Ala Thr Thr  
145 150 155 160

His Met Ile Glu Gln Arg Lys Thr Pro Val Ser Arg Leu Ser Leu Phe  
165 170 175

Ser Phe Lys Gln Ser Pro Pro Gly Thr Pro Val Ser Ser Cys Gln Met  
180 185 190

Arg Phe Asp Gly Pro Leu His Val Pro Met Asn Pro Glu Pro Ala Gly  
195 200 205

Ser His His Val Val Asp Gly Gln Thr Phe Ala Val Pro Asn Pro Ile  
210 215 220

Arg Lys Pro Ala Ser Met Gly Phe Pro Gly Leu Gln Ile Gly His Ala  
225 230 235 240

Ser Gln Leu Leu Asp Thr Gln Val Pro Ser Pro Pro Ser Arg Gly Ser  
245 250 255

Pro Ser Asn Glu Gly Leu Cys Ala Val Cys Gly Asp Asn Ala Ala Cys  
260 265 270

Gln His Tyr Gly Val Arg Thr Cys Glu Gly Cys Lys Gly Phe Phe Lys  
275 280 285

Arg Thr Val Gln Lys Asn Ala Lys Tyr Val Cys Leu Ala Asn Lys Asn  
290 295 300

Cys Pro Val Asp Lys Arg Arg Asn Arg Cys Gln Tyr Cys Arg Phe  
305 310 315 320

Gln Lys Cys Leu Ala Val Gly Met Val Lys Glu Val Val Arg Thr Asp

325	330	335
Ser Leu Lys Gly Arg Arg Gly Arg Leu Pro Ser Lys Pro Lys Ser Pro		
340	345	350
Gln Glu Pro Ser Pro Pro Ser Pro Pro Val Ser Leu Ile Ser Ala Leu		
355	360	365
Val Arg Ala His Val Asp Ser Asn Pro Ala Met Thr Ser Leu Asp Tyr		
370	375	380
Ser Arg Phe Gln Ala Asn Pro Asp Tyr Gln Met Ser Gly Asp Asp Thr		
385	390	395
400		
Gln His Ile Gln Gln Phe Tyr Asp Leu Leu Thr Gly Ser Met Glu Ile		
405	410	415
Ile Arg Gly Trp Ala Glu Lys Ile Pro Gly Phe Ala Asp Leu Pro Lys		
420	425	430
Ala Asp Gln Asp Leu Leu Phe Glu Ser Ala Phe Leu Glu Leu Phe Val		
435	440	445
Leu Arg Leu Ala Tyr Arg Ser Asn Pro Val Glu Gly Lys Leu Ile Phe		
450	455	460
Cys Asn Gly Val Val Leu His Arg Leu Gln Cys Val Arg Gly Phe Gly		
465	470	475
480		
Glu Trp Ile Asp Ser Ile Val Glu Phe Ser Ser Asn Leu Gln Asn Met		
485	490	495
Asn Ile Asp Ile Ser Ala Phe Ser Cys Ile Ala Ala Leu Ala Met Val		
500	505	510
Thr Glu Arg His Gly Leu Lys Glu Pro Lys Arg Val Glu Glu Leu Gln		

515

520

525

Asn Lys Ile Val Asn Cys Leu Lys Asp His Val Thr Phe Asn Asn Gly

530

535

540

Gly Leu Asn Arg Pro Asn Tyr Leu Ser Lys Leu Leu Gly Lys Leu Pro

545

550

555

560

Glu Leu Arg Thr Leu Cys Thr Gln Gly Leu Gln Arg Ile Phe Tyr Leu

565

570

575

Lys Leu Glu Asp Leu Val Pro Pro Pro Ala Ile Ile Asp Lys Leu Phe

580

585

590

Leu Asp Thr Leu Pro Phe

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&lt;211&gt; 22

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

<223> Description of Artificial Sequence:Artificially  
Synthesized Primer Sequence

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22

&lt;210&gt; 6

&lt;211&gt; 22

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

<220>

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Synthesized Primer Sequence

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22

<210> 7

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Artificially  
Synthesized Probe Sequence

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<222> (1)

<223> Label FAM

<220>

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<222> (30)

<223> Label TAMRA

<400> 7

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30

<210> 8

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Artificially  
Synthesized Primer Sequence

<400> 8

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20

<210> 9

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Artificially  
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24

<210> 10

<211> 24

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:Artificially  
Synthesized Probe Sequence

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24

<210> 11

<211> 63

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Artificially  
Synthesized Primer Sequence

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ttt

63

<210> 12

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Artificially  
Synthesized Primer Sequence

<400> 12

tcacccacac tgtgccccatc tacga

25

<210> 13

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Artificially  
Synthesized Primer Sequence

<400> 13

cagcggaacc gctcattgcc aatgg

25

&lt;210&gt; 14

&lt;211&gt; 26

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

<223> Description of Artificial Sequence:Artificially  
Synthesized Probe Sequence

&lt;220&gt;

&lt;221&gt; misc\_binding

&lt;222&gt; (1)

&lt;223&gt; Label FAM

&lt;220&gt;

&lt;221&gt; misc\_binding

&lt;222&gt; (7)

&lt;223&gt; Label TAMRA

&lt;400&gt; 14

atgccctccc ccatgccatc ctgcgt

26